



A

ZNFN3A1 : 1 MEPLKVEKFATANRGNGLRAVTPLRPGELLFRSDPLAYTVCKGSRGVVCDRCLLGKEKL M 60
 ME LKVEKF TANRGNGLRAV PLRPGEELLFRSDPLAYTVCKGSRGVVCDRCLLGKEKL

AK010447: 1 MEALKVEKFTTANRGNGLRAVAPLRPGEELLFRSDPLAYTVCKGSRGVVCDRCLLGKEKL M 60

ZNFN3A1 : 61 RCSQCRVAKYCSAKCOKKAWPDHRECKCLKSCKPRYPPDSVRLLGRVVFKLMDGAPSES 120
 RCSQCR+AKYCSAKCOKKAWPDH+REC CLKSCKPRYPPDSVRLLGRV+ KLMD PSES

AK010447: 61 RCSQCRIAKYCSAKCOKKAWPDHRECSCLKSCKPRYPPDSVRLLGRVIVKLMDEKPSES 120

ZNFN3A1 : 121 EKLYSFYDLESNI+NKLTEDKKEGLRQLVMTFOHFMREEIQQASQLPPAFDLFEAFAKVIC 180
 EKLYSFYDLESNI+NKLTEDKKEGLRQLVMTFOHFMREEIQQASQLPP+FDLFEAFAKVIC

AK010447: 121 EKLYSFYDLESNI+NKLTEDKKEGLRQLAMTFQHFMREEIQQASQLPPSFDLFEAFAKVIC 180

ZNFN3A1 : 181 NSFTTGNAEMOEYGVGLYPSISLNHSCDPNCSTIVFNGPHLLLRAVRDIEVGEELTICYL 240
 NSFTTGNAEMOEYGVGLYPS+SLNHSCDPNCSTIVFNGPHLLLRAVR+IE GEELTICYL

AK010447: 181 NSFTTGNAEMOEYGVGLYPSMSLNHSCDPNCSTIVFNGPHLLLRAVREIEAGEELTICYL 240

ZNFN3A1 : 241 DMLMTSEERRKQLRDQYCFECDCFRCQTQDKADMLTGDEQWKEVQESLKKIEELKAHW 300
 DMLMTSEERRKQLRDQYCFECDC RCQTQDKADMLTGDEQ+WKEVQESLKKIEELKAHW

AK010447: 241 DMLMTSEERRKQLRDQYCFECDCIRCQTQDKADMLTGDEQIWKEVQESLKKIEELKAHW 300

ZNFN3A1 : 301 KWEQVLAMCQAISSNSERLPDINIYQLKVLDCAMDACINLGLLEEALFYGTRTMEPYRI 360
 KWEQVLA+CQAI+ +SNS RLPDINIYQLKVLDCAMDACINLG+LEEALFY RTMEPYRI

AK010447: 301 KWEQVLALCQAIINSNSNRLPDINIYQLKVLDCAMDACINLGMLEEALFYAMRTMEPYRI 360

ZNFN3A1 : 361 FFPGSHPVRCVQVMKVGKLQLHQGMFPQAMKNLRLAFDIMRVTHGREHSLIEDLILLEE 420
 FFPGSHPVRCVQVMKVGKLQLHQGMFPQAMKNLRLAFDIM+VTHGREHSLIEDLILLEE

AK010447: 361 FFPGSHPVRCVQVMKVGKLQLHQGMFPQAMKNLRLAFDIMKVTHGREHSLIEDLILLEE 420

ZNFN3A1 : 421 CDANIRAS 428 (SEQ ID NO: 1)
 CDANIRAS (SEQ ID NO: 81)

AK010447: 421 CDANIRAS 428 (SEQ ID NO: 82)

FIG.2B

BEST AVAILABLE COPY



10/25

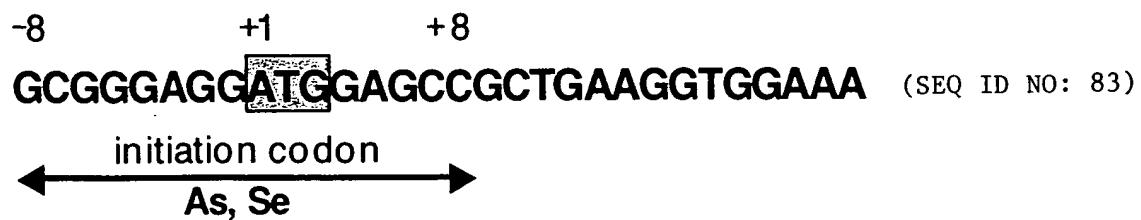


FIG.6A

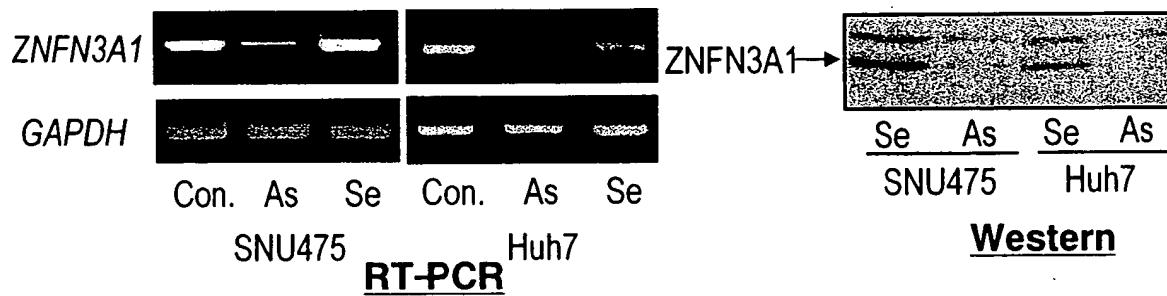


FIG.6B

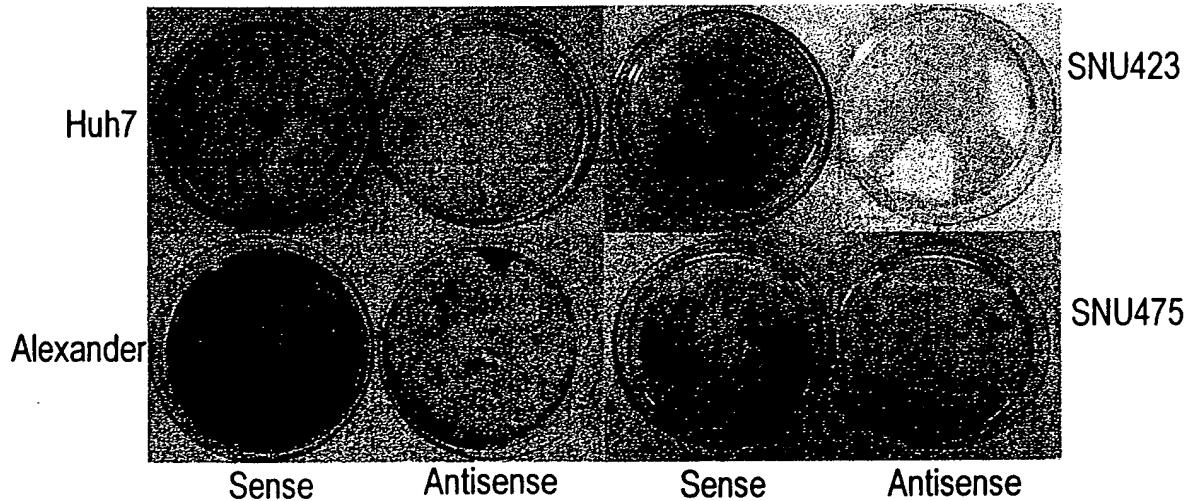


FIG.6C

BEST AVAILABLE COPY